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| BETA. ALPHA. CLUSTER B. CLUSTER B. CLUSTER B. CLUSTER B. | 4370.1; 3. 3. 10. 10. 12. 12. 16; 29; 29; 21. METALLOTHIONEIN_VRT; PARTIAL Bal-thiolate cluster; Chelati | This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bloinformatics and the European Bioinformatics Institute. There are use by non-profit institutions as long as its modified and this statement is not removed. Usage entities requires a license agreement (See http://orrsend an email to license@isb-sib.ch). | FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF FUNCTION: METALLOTHIONEINS HEAVY METALS. RESIDUES THAT BIND VARIOUS HEAVY METALS. DOMAIN: CLASS I METALLOTHIONEINS CONTAIN 2 METAL FOUR DIVALENT TONS ARE CHELATED WITHIN CLUSTER A DOMAIN AND ARE COORDINATED VIA CYSTEINYL THIOLAT CYSTEINE LIGANDS. CLUSTER B, THE CORRESPONDING RECORDINATED VIA CORRESPONDING RESPONDING RECORD THREE DIVALENT IONS TO 9 SIMILARITY: BELONGS TO FAMILY 1 IN METALLOTHIONE SIMILARITY: BELONGS TO FAMILY 1 IN METALLOTHIONE | SEQUENCE FROM N.A. TISSUE=LIVER; MEDLINE=93247066; PubMed=8483164; MEDLINE=93247066; PubMed=8483164; Shartzer K.L., Kage K., Sobieski R.J., Andrews G.K.; Shartzer K.L., Kage K., Sobieski R.J., Andrews G.K.; Evolution of avian metallothionein: DNA sequence ana turkey metallothionein gene and metallothionein cDNAs and quall "; | 23, Created) 23, Last sequence update) 35, Last sequence update) 35, Last annotation update) (MTA) (FRAGMENT). (Common bobwhite). (Chordata; Craniata; Vertebrata; Eut. Neognathae; Galliformes; Phasianidae | PRT; 43 AA | ALIGNMENTS | MT1A_BOVIN MT1A_HORSE MT1A_RABIT MT1B_HORSE MT1B_HORSE MT1B_HORSE MT1B_SHEEP MT1E_SHEEP MT1E_HUMAN MT1E_HUMAN MT1E_HUMAN MT1E_HUMAN MT1G_HUMAN MT1H_BOVIN |
| | PARTIAL. Chelation. | rroduced through a collaboration cics and the EMBL outstation sits are no restrictions on its its content is in no way Usage by and for commercial http://www.isb-sib.ch/announce/ | ENT OF CYSTEINE METAL-BINDING DOMA TER A OF THE ALPHA TIOLATE BRIDGES TO ING REGION WITHIN TO 9 CYSTEINES HIONELN SUPERFAMIL | ndrews G.K.; sequence analyses of the hionein cDNAs from pheasant | re) n bobwhite). rtebrata; Euteleostomi; Phasianidae; | | | P04356 bos taurus P02800 equus cabal P04731 homo sapien P11957 oryctolagus P02801 equus cabal P07438 homo sapien P09577 ovis aries P09578 ovis aries P04733 homo sapien P04733 homo sapien P13640 homo sapien P13640 homo sapien P55942 bos taurus |

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
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FUNCTION: METALLOTHLONEINS HAVE A HIGH CONTENT OF CYSTEINE RESIDUES THAT BIND VARIOUS HEAVY METALS.

DOMAIN: CLASS I METALLOTHLONEINS CONTAIN 2 METAL-BINDING DOMAINS: FOUR DIVALENT IONS ARE CHELATED WITHIN CLUSTER A OF THE ALPHA DOMAIN AND ARE COORDINATED VIA CYSTEINYL THIOLATE BRIDGES TO 11 CYSTEINE LIGANDS. CLUSTER B, THE CORRESPONDING REGION WITHIN THE BETA DOMAIN, CAN LIGATE THREE DIVALENT IONS TO 9 CYSTEINES.

SIMILARITY: BELONGS TO FAMILY 1 IN METALLOTHIONEIN SUPERFAMILY.
                                                                                                                                                                                                                                                                                                                                     European Bioinformatics Institute. The by non-profit institutions as long
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Arch. Biochem. Biochys. 238:18-29(1985).

-i- FUNCTION: THIONINS ARE SMALL PLANT PROTEINS WHICH ARE TOXIC TO ANIMAL CELLS. THEY SEEM TO EXERT THEIR TOXIC EFFECT AT THE LEVEL OF THE CELL MEMBRANE. THE PRECISE FUNCTION, IN PLANTS,
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Vernon L.P., Evett
                                                                                                                                                                                                                                               Pyrularia pubera (Rabbitwood) (Buffalo nut).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Santalales; Santalaceae;
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                                                                                                                                                                                            amino acid sequence."
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P07504;
                                                                                                                  MENDEL; 13438; PYRpu; Thi1; 3. INTERPRO; IPRO01010; -.
                                                                                                                                 HSSP; P01544; 2PLH.
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Pred. No. 43;
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15-DEC-1998
15-DEC-1998
                                                                                                                                        MEDLINE-93011905; PubMed=1397265; Cordeiro M.N., Diniz C.R., Valentim A.C., von E Gilroy J., Richardson M.'.

"The purification and amino acid sequences of from the venom of the Brazilian 'armed' spider
                                                                                                                                                                                                                                    Phoneutria nigriventer (Brazilian armed spider).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Entelegynae; Lycosoidea; Ctenidae; Phoneutria.
                                                                                                                                                                                                                                                                             P29425;
01-APR-1993 (Rel. 25, Created)
01-APR-1993 (Rel. 25, Last sequence update)
01-APR-1993 (Rel. 25, Last annotation updat
                        PIR;
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Kim D.R., Kang K.W.;
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            Venom;
                                   -!- SIMILARITY: TO NEUROTOXINS TX2-1
                                                                                                                    (Keys).";
FEBS Lett.
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                                                      FUNCTION: CAUSES SCRATCHING, LACRIMATION, HYPERSALIVATION, SWEATING AND AGITATION FOLLOWED BY SPASTIC PARALYSIS OF THE ANYERTIOR AND POSTERIOR EXTREMITIES AND DEATH AT DOSE LEVELS 0.79 MG/MOUSE. IT SIGNIFICANTLY ACTIVATES VOLTAGE-DEPENDENT CHANNELS. INSECTICIDAL TO THE LARVAL AND ADULT FORMS OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BE INVOLVED IN LEECH HEMATOPHAGIA SIMILARITY: BELONGS TO THE ANTISTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ino acid sequence of piguamerin, an antistasin-type ibitor from the blood sucking leech Hirudo nipponia.
J. Biochem. 254.692-697(1998).
FUNCTION: INHIBITS PLASMA AND TISSUE KALLIKREIN, AN
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                        S29216; S29216.
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Phoneutria nigriven
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P80302;
01-FEB-1994
01-FEB-1994
15-JUL-1998
                   Hirudo medicinalis (Medicinal leech).
Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida;
Euynchobdellida; Hirudiniformes; Hirudinidae; Hirudo.
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01-APR-1993 (Rel. 25, Created)
01-APR-1993 (Rel. 25, Last sequence up
01-JUL-1993 (Rel. 26, Last annotation
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TX25_PHONI
                                                                                        HIRUSTASIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   spider Phoneutria nigriventer.";
Toxicon 29:1225-1233(1991).
-I-FUNCTION: CAUSES SCRATCHING, LACRIMATION, HYPERSALIVATION,
SWEATING AND AGITATION FOLLOWED BY SPASTIC PARALYSIS OF THE
ANTERIOR AND POSTERIOR EXTREMITIES AND DEATH AT DOSE LEVELS
0.24 MG/MOUSE. INSECTICIDAL TO THE LARVAL AND ADULT FORMS OF
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Entelegynae; Lycosoidea; Ctenidae; Phoneutria.
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-i- SIMILARITY: TO NEUROTOXINS
PIR: B39305; B39305.
PIR: S29215; S29215.
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Cordeiro M.N., Diniz C.R., Valentim A.C., von
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=92196803; PubMed=1801316;
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the 'armed'
                                              Hirudinea;
                                                                                                                                                                                                                                                                                                                                                                                                                                 49
                                                                                                                                                                                                                                                                                                                                                                                    0;
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Best Local :
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MB4C_DROME STANDARD; PRT; 5:

Q01644; Q9VIAO;
Q1-JUL-1993 (Rel. 26, Created)
Q1-JUL-1993 (Rel. 26, Last sequence upda
Q1-CCT-2000 (Rel. 40, Last annotation up
MALE SPECIFIC SPERM PROTEIN MST84DC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Serine protease a ACT_SITE 30 DISULFID 11 DISULFID 21 DISULFID 29 DISULFID 29 DISULFID 33
MEDLINE=20196006; PubMed=10731132;
Adams M.D., Celniker S.E., Holt R.A., Evans C.A.
Amanatides P.G., Scherer S.E., Li P.W., Hoskins
George R.A., Lewis S.E., Richards S., Ashburner
Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q
                                                                                                                                    Kuhn
                                                       STRAIN-BERKELEY;
                                                                  SEQUENCE FROM N.A.
                                                                                                                                           STRAIN=OREGON-R;
MEDLINE=92102953;
                                                                                                                                                                    SEQUENCE FROM
                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Tracheata;
Pterygota; Neoptera; Endopterygota; Diptera
                                                                                                                                                                                                                        Drosophila melanogaster (Fruit fly)
                                                                                                                                                                                                                                     MST84DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Structure
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Sommerhoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mittl P.R., di Marco S., Fendrich G., Pohlig G., Heim J., Sommerhoff C., Fritz H., Priestle J.P., Grutter M.G.; "A new structural class of serine protease inhibitors revealed by structure of the hirustasin-kalllkrein complex.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eur. J. Biochem. 219:937-943(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       serine-proteinase inhibitor from the medical leech Hirudo
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Isolation and characterization of hirustasin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    \overline{\omega}
                                                                                                                                                                                                                                                                                                                                                           29
                                                                                       Drosophila melanogaster."; ch. Dev. 35:143-151(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CHYMOTRYPSIN AND NEUTROPHIL CATHEPSIN G. SIMILARITY: BELONGS TO THE ANTISTASIN FAMILY
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Ef C., Fritz H., Pr
e 5:585-585(1997).
                                                                                                                                   Kuhn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5:253-264(1997).
                                                                                                             of
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                                                                                                                                                                                       eoptera; Endopterygota; Diptera;
Drosophilidae; Drosophila.
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                                                                                                                                   Boersch
                                                                                                                                              PubMed=1684716;
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Priestle
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                                                                                                                                 Glaetzer K.H., Schaefer U.,
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                                 Evans C.A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pohlig
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Grutter M
           Hoskins R.A., Galle R
shburner M., Henderson
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                       Galle R.F
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RESULT 9
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                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                             Matches
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EMBL; AE003672; AAF5403
HSSP; P01180; INPO.
                                                                                                                                                                                                                                                                                                                                                                             Spermatogenesis; Repeat; Multigene family. SEQUENCE 55 AA; 5225 MW; 95A12F3AEC88BD6C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             between the Swiss Institute of Bioinf
the European Bioinformatics Institute
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DOMAIN: THIS PROTEIN IS MOSTLY COMPOSED OF REPETITIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: BELONGS TO THE MST(3)CGP FAMILY.
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                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                   FBgn0004174; Mst84Dc
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                                                                                                                                                                                                                                                                             Conservative
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AAF54025.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Usage
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M87F_DROME STANDA P08175; Q9VFQ6; 01-AUG-1988 (Rel. 08, 01-AUG-1988 (Rel. 08, 01-OCT-2000 (Rel. 40, MALE SPECIFIC SPERM P

. 08, Created)
. 08, Last sequence 40, Last annotations. PROTEIN MST87F

Last sequence up

STANDARD

PRT;

56

AA

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RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams M.D., Celniker S.E., Holt R.A., How, Hoskins R.A., Galle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang O., Chen L.X.,
RA Brandon R.C., Rogers Y.H.C., Blaze's R.G., Champe M., Pfelffer B.D.,
RA Ballaw R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Ballaw R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Glerry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Glodek A., Cong F., Cownes M., Dugan Rocha S., Dunkov B.C., Dunn P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Dup L.E., Downes M., Dugan Rocha S., Dunkov B.C., Dunn P.,
RA Glodek A., Cong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Mattei B., McIntosh T.C., McLeod M.P., McCherson D.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravicz S., Kulp D., Lai Z.,
RA Melson D.R., Nelson K., Nixon K., Nixoskern D.R., Nelson D.L.,
RA Ra Harris N., Moharry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K., Nixon K., Nixoskern D.R., Smith T.,
RA Rollam S.M., Woodage T., Wontey S., Mang A.H., Wang X.,
RA Wang K.H., Land R.S., Pan S., Pollard J., Puri v., Reese M.G.,
RA Wang K.H., Shand R.S., Stapleton M., Strong R., Sun E.,
RA Svirskas R., Tector C., Turner R., Vennter E., Wang A.H., Wang X.,
RA Jehng X.H., Wehs R.F., Zaveri J.S., Zhan M., Zhang G., Zho Q., A.
Ra Jehng X.H., Wens S., Zhan M., Zhang G., Zho Q., A.
Ra Jehng X.H., Wens S., Shan M., Zhang S., Zhu X., Smith 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MST(3)GL-9 (
Drosophila r
EMBL; Y00831; CAA68761
EMBL; AE003702; AAF549
PIR; S00340; WTFF.
                                                                         use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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"Cis-acting regions sufficient for spermatocyte-specific transcriptional and spermatid-specific translational control of the Drosophila melanogaster gene mst(3)gl-9.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=CANTON-S;
MEDLINE=88211557; PubMed=2835228;
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                                                                                                                                                                   European Bioinformatics Institute.
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DEVELOPMENTAL STAGE: PRIMAR
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                                                                                                                                                                                       SWISS-PROT entry is copyright. It is produced through sen the Swiss Institute of Bioinformatics and the EM
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                                                                                                                                                                                                                                                                                                PRIMARY SPERMATOCYTES IS MOSTLY COMPOSED OF
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P46443;
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                       Sparus aurata (Gilthead sed Diesm);
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostoml;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostoi;
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J. Biol. Chem. 270.13879-13884(1995).
-I- FUNCTION: INHIBITS MAMMALIAN ELASTASES
-I- TISSUE SPECIFICITY: NOT FOUND IN THE SI
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Eukaryota; Meta
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FLYBASE; FBgn0002862; Mst87F.
Spermatogenesis; Repeat; Multigene family.
SEQUENCE 56 AA; 5233 MW; 830CD13212C34A47 CRC64;
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Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.
Submitted: Courtion of Cysteine
FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE
RESIDUES THAT BIND VARIOUS HEAVY METALS (BY SIMILARITY).

-!- DOMAIN: CLASS I METALLOTHIONEINS CONTAIN 2 METAL-BINDING DOMAINS.
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Kille P., Olsson P.E.
Submitted (APR-1996)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Zoarcoi
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FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE RESIDUES THAT BIND VARIOUS HEAVY METALS (BY SIMILARITY).

DOMAIN: CLASS I METALLOTHIONEINS CONTAIN 2 METAL-BINDING DOMAINS FOUR DIVALENT IONS ARE CHELATED WITHIN CLUSTER A OF THE ALPHA DOMAIN AND ARE COORDINATED VIA CYSTEINEL THIOLATE BRIDGES TO 11 CYSTEINE LIGANDS. CLUSTER B, THE CORRESPONDING REGION WITHIN THI BETALOTHIONEIN SUPERRAMILY.

SIMILARITY: BELONGS TO FAMILY 1 IN METALLOTHIONEIN SUPERRAMILY.
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METALLOTHIONEIN B (
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Oncorhynchus mykiss (RainDow LLLL,
Oncorhynchus alpinus (Arctic char).
Salvelinus alpinus (Arctic char).
Bukaryota; Metazoa; Chordata; Craniata;
Actinopterygil; Neopterygil; Teleostel;
Actinopterygil; Salmoniformes; Salr
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"Structure of the rainbow trout metallothionein B gene characterization of its metal-responsive region.";
Mol. Cell. Biol. 8:4469-4476(1988).
                                                                                                                                                                              between the Swiss institute. The European Bioinformatics Institute. The European Bioinformatics Institutions as long
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MEDLINE=89039876; PubMed=3185557;
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MEDLINE=88111026; PubMed=2448099;
                                                                                                               or send
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CKKSCCPCCPSGCTKCASGC
                       M18104; AAA49566.1; M22487; AAA49567.1; X59394; CAA42037.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kling P.,
                                                                                                               email to license@isb-sib.ch).
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                                                                                                                                  license agreement (See http://www.isb-sib
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METALLOTHIONEIN B (
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PIR; B27490; B27490.
PIR; S16997; S16997.
PIR; S15509; S15509.
HSSP; P04355; 2MRT.
                                                                                                                                                                                MTB_SALSA
P52720;
01-0CT-1996
             Kille P., Olsson P.E.;

Submitted (APR-1996) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE RESIDUES THAT BIND VARIOUS HEAVY METALS (BY SIMILARITY).

-I- DOMAIN: CLASS I METALLOTHIONEINS CONTAIN 2 METAL-BINDING DOMAINS: FOUR DIVALENT IONS ARE CHELATED WITHIN CLUSTER A OF THE ALPHA DOMAIN AND ARE COORDINATED VIA CYSTEINL THIOLATE BRIDGES TO 11

CYSTEINE LIGANDS. CLUSTER B, THE CORRESPONDING REGION WITHIN THE BETA DOMAIN, CAN LIGATE THREE DIVALENT IONS TO 9 CYSTEINES.

-I- SIMILARITY: BELONGS TO FAMILY I IN METALLOTHIONEIN SUPERFAMILY.
                                                                                                                  Salmo salar (Atlantic salmon).
Salmo salar (Atlantic salmon).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
Acthopterygii; Meopterygii; Teleostei; Buteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Salmo.
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PROSITE; PS00203; METALLOTHIONEIN_VRT;
Metal-binding; Metal-thiolate cluster;
                                                                                            TISSUE=LIVER;
                                                                                                   SEQUENCE FROM N.A.
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INTERPRO; IPRO
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entities requires a license agreement 'Com'
                metallothionein, AmMT-A, and expression of the gene dur development.";
Submitted (JUN-1997) to the EMBL/GenBank/DDBJ databases
                                                               Saint-Jacques
                                                                           TISSUE=EMBRYO;
                                                                                                              Eukaryota;
Amphibia; I
                                                                                                                         Ambystoma mexicanum (Axolotl).
Eukaryota; Metazoa; Chordata;
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187A9D0FBE5BD967 CRC64;
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                                                                                                              Craniata; Vertebrata; Euteleostomi; Salamandroidea; Ambystomatidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS00203; METALLOTHIONEIN_VRT;
Metal-binding; Metal-thiolate cluster;
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                                   1 CXXXCXXXXXXXXXXXXXXXXX
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